



SEQUENCE LISTING

<110> Dragic, Tatjana
Olson, William C.

<120> SULFATED CCR5 PEPTIDES FOR HIV-1 INFECTION

<130> 61010-AB-1

<140> US 10/086,814

<141> 2002-02-28

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<170> PatentIn version 3.1

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Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
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more than 2 amino acids, they have a sequence identical to the sequence set forth in SEQ ID NO: 1 beginning with the Ile at position 9 and extending therefrom in the amino terminal direction.

<220>
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 <222> (7)..(8)
 <223> SULFATATION

<220>
 <221> MISC_FEATURE
 <222> (11)..(11)
 <223> Xaa represents from 0 to 334 amino acids, where if there are more than 2 amino acids, they have a sequence identical to the sequence set forth in SEQ ID NO: 1 beginning with the Glu at position 18 and extending therefrom in the carboxy terminal direction.

<220>
 <221> MISC_FEATURE
 <222> (12)..(12)
 <223> Xaa is any amino acid

<400> 19

Xaa Xaa Tyr Asp Ile Asn Tyr Tyr Thr Ser Xaa Xaa
 1 5 10

<210> 20
 <211> 12
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (1)..(1)
 <223> Xaa is any amino acid

<220>
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 <223> ACETYLTATION

<220>
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 <223> Xaa represents from 0 to 9 amino acids, where if there are
 more than 2 amino acids, they have a sequence identical to the se-
 quence set forth in SEQ ID NO: 1 beginning with the Ile at posi-
 tion 9 and extending therefrom in the amino terminal direction.

<220>
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 <223> SULFATATION

<220>
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 <222> (8)..(8)
 <223> SULFATATION

<220>
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 <222> (11)..(11)
 <223> Xaa represents from 0 to 334 amino acids, where if there are
 more than 2 amino acids, they have a sequence identical to the

sequen

ce set forth in SEQ ID NO: 1 beginning with the Glu at posi
tion 1
8 and extending therefrom in the carboxy terminal direction
.

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa is any amino acid

<400> 20

Xaa	Xaa	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser	Xaa	Xaa
1				5					10		

<210> 21

<211> 12

<212> PRT

<213> Homo sapiens

<220>

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<222> (1)..(1)

<223> Xaa is any amino acid

<220>

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<222> (2)..(2)

<223> Xaa represents from 0 to 9 amino acids, where if there are
more t

han 2 amino acids, they have a sequence identical to the se
quence

set forth in SEQ ID NO: 1 beginning with the Ile at positi
on 9 a

nd extending therefrom in the amino terminal direction.

<220>

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 <222> (3)..(3)
 <223> SULFATATION

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 <223> SULFATATION

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 <223> Xaa represents from 0 to 334 amino acids, where if there are more than 2 amino acids, they have a sequence identical to the sequence set forth in SEQ ID NO: 1 beginning with the Glu at position 18 and extending therefrom in the carboxy terminal direction.

<220>
 <221> MISC_FEATURE
 <222> (12)..(12)
 <223> Xaa is any amino acid

<220>
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 <222> (12)..(12)
 <223> AMIDATION

<400> 21

Xaa Xaa Tyr Asp Ile Asn Tyr Tyr Thr Ser Xaa Xaa
 1 5 10

<210> 22

<211> 12
 <212> PRT
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 <223> Xaa is any amino acid

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<220>
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 <223> SULFATATION

<220>
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<220>
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<222> (12)..(12)
 <223> Xaa is any amino acid

<220>
 <221> MOD_RES
 <222> (12)..(12)
 <223> AMIDATION

<400> 22

Xaa	Xaa	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser	Xaa	Xaa
1				5					10		

<210> 23
 <211> 12
 <212> PRT
 <213> Homo sapiens

<220>
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 <223> Xaa is any amino acid

<220>
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 <222> (2)..(2)
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 more than 2 amino acids, they have a sequence identical to the se
 quence set forth in SEQ ID NO: 1 beginning with the Ile at positi
 on 9 and extending therefrom in the amino terminal direction.

<220>
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 <223> SULFATATION

<220>
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 <223> SULFATATION

<220>
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<220>
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 <222> (12)..(12)
 <223> Xaa is any amino acid

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<400> 23

Xaa Xaa Tyr Asp Ile Asn Tyr Tyr Thr Ser Xaa Xaa
 1 5 10

<210> 24
 <211> 12
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 <223> Xaa is any amino acid

<220>
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 <223> ACETYLTATION

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 on 9 and extending therefrom in the amino terminal direction.

<220>
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 <223> SULFATATION

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 <223> Xaa represents from 0 to 334 amino acids, where if there ar
 e more than 2 amino acids, they have a sequence identical to the

sequen

ce set forth in SEQ ID NO: 1 beginning with the Glu at position 1
8 and extending therefrom in the carboxy terminal direction

<220>

<221> MISC_FEATURE

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<223> Xaa is any amino acid

<220>

<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 24

Xaa	Xaa	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser	Xaa	Xaa
1				5					10		

<210> 25

<211> 12

<212> PRT

<213> Homo sapiens

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<223> Xaa is any amino acid

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<222> (1)..(1)

<223> ACETYLATION

<220>

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 on 9 and extending therefrom in the amino terminal direction.

<220>
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 <222> (7)..(8)
 <223> SULFATATION

<220>
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 <223> Xaa represents from 0 to 334 amino acids, where if there ar
 e more than 2 amino acids, they have a sequence identical to the
 sequen ce set forth in SEQ ID NO: 1 beginning with the Glu at posi
 tion 1 8 and extending therefrom in the carboxy terminal direction

<220>
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 <222> (12)..(12)
 <223> Xaa is any amino acid

<220>
 <221> MOD_RES
 <222> (12)..(12)
 <223> AMIDATION

<400> 25

Xaa Xaa Tyr Asp Ile Asn Tyr Tyr Thr Ser Xaa Xaa
1 5 10

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<210> 26
<211> 12
<212> PRT
<213> Homo sapiens
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<222>  (1)..(1)
<223>  Xaa is any amino acid
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<220>
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<222> (1) .. (1)
<223> ACETYLATION
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<220>
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<223>  Xaa represents from 0 to 9 amino acids, where if there are
more t      han 2 amino acids, they have a sequence identical to the se
quence      set forth in SEQ ID NO: 1 beginning with the Ile at positi
on 9 a      nd extending therefrom in the amino terminal direction.

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<220>
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<223> SULFATATION

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<222> (8) . (8)
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<223> SULFATATION

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents from 0 to 334 amino acids, where if there are more

than 2 amino acids, they have a sequence identical to the sequence set forth in SEQ ID NO: 1 beginning with the Glu at position 1

8 and extending therefrom in the carboxy terminal direction

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa is any amino acid

<220>

<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 26

Xaa	Xaa	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser	Xaa	Xaa
1				5					10		

<210> 27

<211> 17

<212> PRT

<213> Homo sapiens

<400> 27

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 28
 <211> 17
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (2)..(2)
 <223> SULFATATION

<220>
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 <222> (9)..(9)
 <223> SULFATATION

<220>
 <221> MOD_RES
 <222> (13)..(13)
 <223> SULFATATION

<400> 28

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 29
 <211> 17
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (9) .. (9)
 <223> SULFATATION

<220>
 <221> MOD_RES
 <222> (13) .. (13)
 <223> SULFATATION

<400> 29

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 30
 <211> 21
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (9) .. (9)
 <223> SULFATATION

<220>
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 <222> (13) .. (13)
 <223> SULFATATION

<220>
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 <222> (21) .. (21)

<223> BIOTIN

<400> 30

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu	Gly	Ala	Gly	Lys
			20	

<210> 31

<211> 17

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (9)..(9)

<223> SULFATATION

<400> 31

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 32

<211> 17

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (13)..(13)

<223> SULFATATION

<400> 32

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 33
 <211> 5
 <212> PRT
 <213> Homo sapiens

<220>
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 <223> SULFATATION

<220>
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 <222> (5)..(5)
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<400> 33

Tyr	Asp	Ile	Asn	Tyr
1			5	

<210> 34
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 <212> PRT
 <213> Homo sapiens

<220>
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<223> PHOSPHORYLATION

<220>

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<222> (9)..(9)

<223> PHOSPHORYLATION

<220>

<221> MOD_RES

<222> (13)..(13)

<223> PHOSPHORYLATION

<400> 34

Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	Ser
1				5					10					15	

Glu

<210> 35

<211> 17

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (9)..(9)

<223> PHOSPHORYLATION

<220>

<221> MOD_RES

<222> (13)..(13)

<223> PHOSPHORYLATION

<400> 35

Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr Ser
 1 5 10 15

Glu

<210> 36
 <211> 21
 <212> PRT
 <213> Homo sapiens

<220>
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 <223> PHOSPHORYLATION

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 <223> PHOSPHORYLATION

<220>
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 <222> (21)..(21)
 <223> BIOTIN

<400> 36

Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr Ser
 1 5 10 15

Glu Gly Ala Gly Lys
 20

<210> 37

<211> 17
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (1)..(1)
 <223> SULFATATION

<220>
 <221> MOD_RES
 <222> (11)..(11)
 <223> SULFATATION

<400> 37

Tyr	Val	Ser	Gln	Pro	Asp	Asn	Thr	Tyr	Ile	Tyr	Ser	Tyr	Glu	Ser	Ile
1				5					10					15	

Asp

<210> 38
 <211> 17
 <212> PRT
 <213> Homo sapiens

<220>
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 <222> (9)..(9)
 <223> SULFATATION

<220>
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 <222> (13)..(13)
 <223> SULFATATION

<400> 38

Ser	Ile	Asp	Ile	Tyr	Asn	Pro	Thr	Tyr	Val	Ser	Asn	Tyr	Glu	Ser	Asp
1				5					10					15	

Tyr